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SEQUENCE LISTING

<110> Langley, Ries
Fraser, John David

<120> Set 1 Proteins And Uses Thereof

<130> 55502-012us1

<140> 10/594,291

<141> 2004-12-07

<150> AU 2004901570

<151> 2004-03-24

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<170> PatentIn version 3.5

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Lys Gln Glu Arg Val Gln His Leu His Asp Ile Arg Asp Leu His Arg
35 40 45

Tyr Tyr Ser Ser Glu Ser Phe Glu Tyr Ser Asn Val Ser Gly Lys Val
50 55 60

Glu Asn Tyr Asn Gly Ser Asn Val Val Arg Phe Asn Pro Lys Asp Gln
65 70 75 80

Asn His Gln Leu Phe Leu Leu Gly Lys Asp Lys Glu Gln Tyr Lys Glu
85 90 95

Gly Leu Gln Gly Gln Asn Val Phe Val Val Gln Glu Leu Ile Asp Pro
100 105 110

Asn Gly Arg Leu Ser Thr Val Gly Gly Val Thr Lys Lys Asn Asn Lys
115 120 125

Thr Ser Glu Thr Asn Thr Pro Leu Phe Val Asn Lys Val Asn Gly Glu
130 135 140

Asp Leu Asp Ala Ser Ile Asp Ser Phe Leu Ile Gln Lys Glu Glu Ile
Page 1

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Asp Leu Asp Ala Ser Ile Asp Ser Phe Leu Ile Gln Lys Glu Glu Ile
145 150 155 160

Ser Leu Lys Glu Leu Asp Phe Lys Ile Arg Gln Gln Leu Val Asn Asn
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Tyr Gly Leu Tyr Lys Gly Thr Ser Lys Tyr Gly Lys Ile Ile Ile Asn
 180 185 190

Leu Lys Asp Glu Asn Lys Val Glu Ile Asp Leu Gly Asp Lys Leu Gln
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Ser Val Thr Ile Asn Gln Ile
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Thr Thr Gly Val Ile Thr Ser Glu Gly Gln Ala Val His Ala Lys Glu
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Lys Gln Glu Arg Val Gln His Leu Tyr Asp Ile Lys Asp Leu Tyr Arg
35 40 45

Tyr Tyr Ser Ser Glu Ser Phe Glu Phe Ser Asn Ile Ser Gly Lys Val
50 55 60

Glu Asn Tyr Asn Gly Ser Asn Val Val Arg Phe Asn Gln Glu Lys Gln
65 70 75 80

Asn His Gln Leu Phe Leu Leu Gly Lys Asp Lys Asp Lys Tyr Lys Lys
85 90 95

Gly Leu Glu Gly Gln Asn Val Phe Val Val Lys Glu Leu Ile Asp Pro
100 105 110

Asn Gly Arg Leu Ser Thr Val Gly Gly Val Thr Lys Lys Asn Asn Lys
 Page 3

115					120					sequence					125				
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Tyr	Gly	Leu	Tyr	Lys	Gly	Thr	Thr	Lys	Tyr	Gly	Lys	Ile	Thr	Ile	Asn				
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Leu	Lys	Asp	Glu	Lys	Lys	Glu	Val	Ile	Asp	Leu	Gly	Asp	Lys	Leu	Gln				
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			20					25					30						
Lys	Gln	Glu	Arg	Val	Gln	Glu	Leu	Tyr	Asp	Ile	Lys	Asp	Leu	Tyr	Arg				
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Lys Val Glu Asn Tyr Asn Gly Ser Asn Val Val Arg Phe Asn Gln Asp
35 40 45

Gly Gln Asn His Gln Leu Phe Leu Leu Gly Glu Asp Lys Ala Lys Tyr
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Lys Glu Gly Leu Glu Gly Glu Asn Val Phe Val Val Lys Glu Leu Ile

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 35 40 45
 Lys Gln Asn His Gln Leu Phe Leu Leu Gly Glu Asp Lys Ala Lys Tyr
 50 55 60
 Lys Gln Gly Leu Gln Gly Gln Asp Val Phe Val Val Lys Glu Leu Ile
 65 70 75 80
 Asp Pro Asn Gly Arg Leu Ser Thr Val Gly Gly Val Thr Lys Lys Asn

sequence																	
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	aaagctaaat acaaacaggg tctgcagggt caggacgttt tcgttgtaa agaactgac	240
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 35 40 45

Asn Gln Asn His Gln Leu Phe Leu Ser Gly Lys Asp Lys Asp Lys Tyr
 50 55 60

Lys Glu Gly Leu Glu Gly Gln Asn Val Phe Val Val Lys Glu Leu Ile
 65 70 75 80

Asp Pro Asn Gly Arg Leu Ser Thr Val Gly Gly Val Thr Lys Lys Asn
 85 90 95

Asn Gln Ser Ser Glu Thr Asn Thr Pro Leu Phe Ile Lys Lys Val Tyr
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Gly Gly Asn Leu Asp Ala Ser Ile Glu Ser Phe Leu Ile Asn Lys Glu
 115 120 125

sequence

Glu Val Ser Leu Lys Glu Leu Asp Phe Lys Ile Arg Gln His Leu Val
130 135 140

Lys Asn Tyr Gly Leu Tyr Lys Gly Thr Thr Lys Tyr Gly Lys Ile Thr
145 150 155 160

Phe Asn Leu Lys Asp Gly Glu Lys Gln Glu Ile Asp Leu Gly Asp Lys
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35 40 45

Asn Gln Asn His Gln Leu Phe Leu Leu Gly Lys Asp Lys Glu Lys Tyr
50 55 60

Lys Glu Gly Ile Glu Gly Lys Asp Val Phe Val Val Lys Glu Leu Ile
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Asp Pro Asn Gly Arg Leu Ser Thr Val Gly Gly Val Thr Lys Lys Asn
85 90 95

Asn Lys Ser Ser Glu Thr Asn Thr His Leu Phe Val Asn Lys Val Tyr
100 105 110

Gly Gly Asn Leu Asp Ala Ser Ile Asp Ser Phe Ser Ile Asn Lys Glu
115 120 125

Glu Val Ser Leu Lys Glu Leu Asp Phe Lys Ile Arg Gln His Leu Val
130 135 140

sequence

Lys Asn Tyr Gly Leu Tyr Lys Gly Thr Thr Lys Tyr Gly Lys Ile Thr
145 150 155 160

Ile Asn Leu Lys Asp Gly Glu Lys Gln Glu Ile Asp Leu Gly Asp Lys
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35 40 45

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50 55 60

Lys Lys Gly Leu Glu Gly Gln Asn Val Phe Val Val Lys Glu Leu Ile
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Asp Pro Asn Gly Arg Leu Ser Thr Val Gly Gly Val Thr Lys Lys Asn
85 90 95

Asn Lys Ser Ser Glu Thr Asn Thr His Leu Phe Val Asn Lys Val Tyr
100 105 110

Gly Gly Asn Leu Asp Ala Ser Ile Asp Ser Phe Leu Ile Asn Lys Glu
115 120 125

Glu Val Ser Leu Lys Glu Leu Asp Phe Lys Ile Arg Lys Gln Leu Val
130 135 140

Glu Lys Tyr Gly Leu Tyr Lys Gly Thr Thr Lys Tyr Gly Lys Ile Thr
145 150 155 160

sequence

Ile Asn Leu Lys Asp Glu Lys Lys Glu Val Ile Asp Leu Gly Asp Lys
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35 40 45

Lys Gln Asn His Gln Leu Phe Leu Leu Gly Lys Asp Lys Asp Lys Tyr
50 55 60

Lys Lys Gly Leu Glu Gly Gln Asn Val Phe Val Val Lys Glu Leu Ile
65 70 75 80

Asp Pro Asn Gly Arg Leu Ser Thr Val Gly Gly Val Thr Lys Lys Asn
85 90 95

Asn Lys Ser Ser Glu Thr Asn Thr His Leu Phe Val Asn Lys Val Tyr
100 105 110

Gly Gly Asn Leu Asp Ala Ser Ile Asp Ser Phe Leu Ile Asn Lys Glu
115 120 125

Glu Val Ser Leu Lys Glu Leu Asp Phe Lys Ile Arg Lys Gln Leu Val
130 135 140

Glu Lys Tyr Gly Leu Tyr Lys Gly Thr Thr Lys Tyr Gly Lys Ile Thr
145 150 155 160

Ile Asn Leu Lys Asp Glu Lys Lys Glu Val Ile Asp Leu Gly Asp Lys
165 170 175

sequence

Leu Gln Phe Glu Arg Met Gly Asp Val Leu Asn Ser Lys Asp Ile Gln
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Asn Ile Ala Val Thr Ile Asn Gln Ile
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Lys Val Glu Asn Tyr Asn Gly Ser Asn Val Val Arg Phe Asn Pro Lys
35 40 45

Asp Gln Asn His Gln Leu Phe Leu Leu Gly Lys Asp Lys Glu Gln Tyr
50 55 60

Lys Glu Gly Leu Gln Gly Gln Asn Val Phe Val Val Gln Glu Leu Ile
65 70 75 80

Asp Pro Asn Gly Arg Leu Ser Thr Val Gly Gly Val Thr Lys Lys Asn
85 90 95

Asn Lys Thr Ser Glu Thr Asn Thr Pro Leu Phe Val Asn Lys Val Asn
100 105 110

Gly Glu Asp Leu Asp Ala Ser Ile Asp Ser Phe Leu Ile Gln Lys Glu
115 120 125

Glu Ile Ser Leu Lys Glu Leu Asp Phe Lys Ile Arg Gln Gln Leu Val
130 135 140

Asn Asn Tyr Gly Leu Tyr Lys Gly Thr Ser Lys Tyr Gly Lys Ile Ile
145 150 155 160

Ile Asn Leu Lys Asp Glu Asn Lys Val Glu Ile Asp Leu Gly Asp Lys
165 170 175

Leu Gln Phe Glu Arg Met Gly Asp Val Leu Asn Ser Lys Asp Ile Arg
180 185 190

sequence
Gly Ile Ser Val Thr Ile Asn Gln Ile
195 200